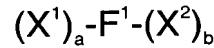


What is claimed is:

1. A composition of matter of the formula



and multimers thereof, wherein:

5 F¹ is an Fc domain;

X¹ and X² are each independently selected from -(L¹)_c-P¹, -(L¹)_c-P¹-(L²)_d-P², -(L¹)_c-P¹-(L²)_d-P²-(L³)_e-P³, and -(L¹)_c-P¹-(L²)_d-P²-(L³)_e-P³-(L⁴)_f-P⁴

10 P¹, P², P³, and P⁴ are each independently sequences of pharmacologically active peptides;

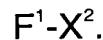
L¹, L², L³, and L⁴ are each independently linkers; and

a, b, c, d, e, and f are each independently 0 or 1, provided that at least one of a and b is 1.

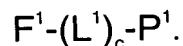
2. The composition of matter of Claim 1 of the formulae



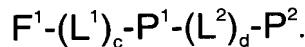
15 or



3. The composition of matter of Claim 1 of the formula



- 20 4. The composition of matter of Claim 1 of the formula



5. The composition of matter of Claim 1 wherein F¹ is an IgG Fc domain.

6. The composition of matter of Claim 1 wherein F¹ is an IgG1 Fc domain.

- 25 7. The composition of matter of Claim 1 wherein F¹ comprises the sequence of SEQ ID NO: 2.

8. The composition of matter of Claim 1 wherein X¹ and X² comprise an IL-1 antagonist peptide sequence.

9. The composition of matter of Claim 8 wherein the IL-1 antagonist peptide sequence is selected from SEQ ID NOS: 212, 907, 908, 909, 910, 917, and 979.
10. The composition of matter of Claim 8 wherein the IL-1 antagonist peptide sequence is selected from SEQ ID NOS: 213 to 271, 671 to 906, 911 to 916, and 918 to 1023.
11. The composition of matter of Claim 8 wherein F¹ comprises the sequence of SEQ ID NO: 2.
12. The composition of matter of Claim 1 wherein X¹ and X² comprise an EPO-mimetic peptide sequence.
13. The composition of matter of Claim 12 wherein the EPO-mimetic peptide sequence is selected from Table 5.
14. The composition of matter of Claim 12 wherein F¹ comprises the sequence of SEQ ID NO: 2.
15. The composition of matter of Claim 12 comprising a sequence selected from SEQ ID NOS: 83, 84, 85, 124, 419, 420, 421, and 461..
16. The composition of matter of claim 12 comprising a sequence selected from SEQ ID NOS: 339 and 340.
17. The composition of matter of Claim 12 comprising a sequence selected from SEQ ID NOS: 20 and 22.
18. The composition of matter of Claim 3 wherein P¹ is a TPO-mimetic peptide sequence.
19. The composition of matter of Claim 18 wherein P¹ is a TPO-mimetic peptide sequence selected from Table 6.
20. The composition of matter of Claim 18 wherein F¹ comprises the sequence of SEQ ID NO: 2.
21. The composition of matter of Claim 18 having a sequence selected from SEQ ID NOS: 6 and 12.
22. A DNA encoding a composition of matter of any of Claims 1 to 21.

23. An expression vector comprising the DNA of Claim 22.
24. A host cell comprising the expression vector of Claim 23.
25. The cell of Claim 24, wherein the cell is an E. coli cell.
26. A process for preparing a pharmacologically active compound,
5 which comprises
 - a) selecting at least one randomized peptide that modulates the activity of a protein of interest; and
 - b) preparing a pharmacologic agent comprising at least one Fc domain covalently linked to at least one amino acid sequence of the selected peptide or peptides.
- 10 27. The process of Claim 26, wherein the peptide is selected in a process comprising screening of a phage display library, an E. coli display library, a ribosomal library, or a chemical peptide library.
28. The process of Claim 26, wherein the preparation of the
15 pharmacologic agent is carried out by:
 - a) preparing a gene construct comprising a nucleic acid sequence encoding the selected peptide and a nucleic acid sequence encoding an Fc domain; and
 - b) expressing the gene construct.
- 20 29. The process of Claim 26, wherein the gene construct is expressed in an E. coli cell.
30. The process of Claim 26, wherein the protein of interest is a cell surface receptor.
31. The process of Claim 26, wherein the protein of interest has a linear
25 epitope.
32. The process of Claim 26, wherein the protein of interest is a cytokine receptor.
33. The process of Claim 26, wherein the peptide is an EPO-mimetic peptide.

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34. The process of Claim 26, wherein the peptide is a TPO-mimetic peptide.
35. The process of Claim 26, wherein the peptide is an IL-1 antagonist peptide.
- 5 36. The process of Claim 26, wherein the peptide is an MMP inhibitor peptide or a VEGF antagonist peptide.
37. The process of Claim 26, wherein the peptide is a TNF-antagonist peptide.
38. The process of Claim 26, wherein the peptide is a CTLA4-mimetic peptide.
- 10 39. The process of Claim 26, wherein the peptide is selected from Tables 4 to 20.
40. The process of Claim 26, wherein the selection of the peptide is carried out by a process comprising:
 - 15 a) preparing a gene construct comprising a nucleic acid sequence encoding a first selected peptide and a nucleic acid sequence encoding an Fc domain;
 - b) conducting a polymerase chain reaction using the gene construct and mutagenic primers, wherein
 - 20 i) a first mutagenic primer comprises a nucleic acid sequence complementary to a sequence at or near the 5' end of a coding strand of the gene construct, and
 - ii) a second mutagenic primer comprises a nucleic acid sequence complementary to the 3' end of the noncoding strand of the gene construct.
- 25 41. The process of Claim 26, wherein the compound is derivatized.
42. The process of Claim 26, wherein the derivatized compound comprises a cyclic portion, a cross-linking site, a non-peptidyl

linkage, an N-terminal replacement, a C-terminal replacement, or a modified amino acid moiety.

43. The process of Claim 26 wherein the Fc domain is an IgG Fc domain.
- 5 44. The process of Claim 26, wherein the vehicle is an IgG1 Fc domain.
45. The process of Claim 26, wherein the vehicle comprises the sequence of SEQ ID NO: 2.
46. The process of Claim 26, wherein the compound prepared is of the formula

10
$$(X^1)_a - F^1 - (X^2)_b$$

and multimers thereof, wherein:

F^1 is an Fc domain;

X^1 and X^2 are each independently selected from $-(L^1)_c - P^1$, $-(L^1)_c - P^1 - (L^2)_d - P^2$, $-(L^1)_c - P^1 - (L^2)_d - P^2 - (L^3)_e - P^3$, and $-(L^1)_c - P^1 - (L^2)_d - P^2 - (L^3)_e - P^3 - (L^4)_f - P^4$

P^1 , P^2 , P^3 , and P^4 are each independently sequences of pharmacologically active peptides;

L^1 , L^2 , L^3 , and L^4 are each independently linkers; and

a, b, c, d, e, and f are each independently 0 or 1, provided

20 that at least one of a and b is 1.

47. The process of Claim 46, wherein the compound prepared is of the formulae

$$X^1 - F^1$$

or

$$F^1 - X^2.$$

- 25 48. The process of Claim 46, wherein the compound prepared is of the formulae

$$F^1 - (L^1)_c - P^1$$

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or

$$F^1-(L^1)_c-P^1-(L^2)_d-P^2.$$

49. The process of Claim 46, wherein F¹ is an IgG Fc domain.
50. The process of Claim 46, wherein F¹ is an IgG1 Fc domain.
- 5 51. The process of Claim 46, wherein F¹ comprises the sequence of SEQ ID NO: 2.